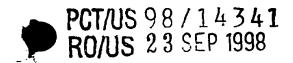


## SEQUENCE LISTING

The nucleotide sequence (SEQ ID NO:1) and deduced amino acid sequence (SEQ ID NO:2) of *Pseudomonas* exotoxin A are:

	5	GCC Ala 1	GAA Glu	GAA Glu	GCT Ala	TTC Phe 5	GAC Asp	CTC Leu	TGG Trp	AAC Asn	GAA Glu 10	TGC Cys	GCC Ala	AAA Lys	GCC Ala	TGC Cys 15	GTG Val	41	3
	10	CTC Leu	GAC Asp	CTC Leu	AAG Lys 20	GAC Asp	GGC Gly	GTG Val	CGT Arg	TCC Ser 25	AGC Ser	CGC Arg	ATG Met	AGC Ser	GTC Val 30	GAC Asp	CCG Pro	90	5
	15	GCC Ala	ATC Ile	GCC Ala 35	GAC Asp	ACC Thr	AAC Asn	GGC Gly	CAG Gln 40	GGC Gly	GTG Val	CTG Leu	CAC His	TAC Tyr 45	TCC Ser	ATG Met	GTC Val	144	4
		CTG Leu	GAG Glu 50	GGC Gly	GGC Gly	AAC Asn	GAC Asp	GCG Ala 55	CTC Leu	AAG Lys	CTG Leu	GCC Ala	ATC Ile 60	GAC Asp	AAC Asn	GCC Ala	CTC Leu	19:	2
T	20	AGC Ser 65	ATC Ile	ACC Thr	AGC Ser	GAC Asp	GGC Gly 70	CTG Leu	ACC Thr	ATC Ile	CGC Arg	CTC Leu 75	GAA Glu	GGC Gly	GGC Gly	GTC Val	GAG Glu 80	24	0
	25	CCG Pro	AAC Asn	AAG Lys	CCG Pro	GTG Val 85	CGC Arg	TAC Tyr	AGC Ser	TAC Tyr	ACG Thr 90	CGC Arg	CAG Gln	GCG Ala	CGC Arg	GGC Gly 95	AGT Ser	28	В
	30	TGG Trp	TCG Ser	CTG Leu	AAC Asn 100	TGG Trp	CTG Leu	GTA Val	CCG Pro	ATC Ile 105	GGC Gly	CAC His	GAG Glu	AAG Lys	CCC Pro 110	TCG Ser	AAC Asn	330	5
	35	ATC Ile	AAG Lys	GTG Val 115	TTC Phe	ATC Ile	CAC His	GAA Glu	CTG Leu 120	AAC Asn	GCC Ala	GGC Gly	AAC Asn	CAG Gln 125	CTC Leu	AGC Ser	CAC His	384	4
	40	ATG Met	TCG Ser 130	CCG Pro	ATC Ile	TAC Tyr	ACC Thr	ATC Ile 135	GAG Glu	ATG Met	GGC Gly	GAC Asp	GAG Glu 140	TTG Leu	CTG Leu	GCG Ala	AAG Lys	43:	2
	40	CTG Leu 145	GCG Ala	CGC Arg	GAT Asp	GCC Ala	ACC Thr 150	TTC Phe	TTC Phe	GTC Val	AGG Arg	GCG Ala 155	CAC His	GAG Glu	AGC Ser	AAC Asn	GAG Glu 160	480	O
	45	ATG Met	CAG Gln	CCG Pro	ACG Thr	CTC Leu 165	GCC Ala	ATC Ile	AGC Ser	CAT His	GCC Ala 170	GGG Gly	GTC Val	AGC Ser	GTG Val	GTC Val 175	ATG Met	528	3
	50	GCC Ala	CAG Gln	ACC Thr	CAG Gln 180	CCG Pro	CGC Arg	CGG Arg	GAA Glu	AAG Lys 185	CGC Arg	TGG Trp	AGC Ser	GAA Glu	TGG Trp 190	GCC Ala	AGC Ser	576	5
	55	GGC Gly	AAG Lys	GTG Val 195	TTG Leu	TGC Cys	CTG Leu	CTC Leu	GAC Asp 200	CCG Pro	CTG Leu	GAC Asp	GGG Gly	GTC Val 205	TAC Tyr	AAC Asn	TAC Tyr	624	1
	۲۸	CTC Leu	GCC Ala 210	CAG Gln	CAA Gln	CGC Arg	TGC Cys	AAC Asn 215	CTC Leu	GAC Asp	GAT Asp	ACC Thr	TGG Trp 220	GAA Glu	GGC Gly	AAG Lys	ATC Ile	672	2
•	60	TAC Tyr 225	CGG Arg	GTG Val	CTC Leu	GCC Ala	GGC Gly 230	AAC Asn	CCG Pro	GCG Ala	AAG Lys	CAT His 235	GAC Asp	CTG Leu	GAC Asp	ATC Ile	AAA Lys 240	720	)





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,		CCC Pro	ACG Thr	GTC Val	ATC Ile	AGT Ser 245	CAT His	CGC Arg	CTG Leu	CAC His	TTT Phe 250	CCC Pro	GAG Glu	GGC Gly	GGC Gly	AGC Ser 255	CTG Leu	768
	5	GCC Ala	GCG Ala	CTG Leu	ACC Thr 260	GCG Ala	CAC His	CAG Gln	GCT Ala	TGC Cys 265	CAC His	CTG Leu	CCG Pro	CTG Leu	GAG Glu 270	ACT Thr	TTC Phe	816
	10	ACC Thr	CGT Arg	CAT His 275	CGC Arg	CAG Gln	CCG Pro	CGC Arg	GGC Gly 280	TGG Trp	GAA Glu	CAA Gln	CTG Leu	GAG Glu 285	CAG Gln	TGC Cys	GGC Gly	864
	15	Tyr	Pro 290	Val	Gln	Arg	Leu	Val 295	Ala	Leu	Tyr	Leu	300	Ala	CGG Arg	Leu	ser	912
	20	Trp 305	Asn	Gln	Val	Asp	Gln 310	Val	Ile	Arg	Asn	Ala 315	Leu	Ala	AGC Ser	Pro	320	960
<u> </u>	25	AGC Ser	GGC Gly	GGC Gly	GAC Asp	CTG Leu 325	GGC Gly	GAA Glu	GCG Ala	ATC Ile	CGC Arg 330	GAG Glu	CAG Gln	CCG Pro	GAG Glu	CAG Gln 335	GCC Ala	1008
Ó	25	CGT Arg	CTG Leu	GCC Ala	CTG Leu 340	ACC Thr	CTG Leu	GCC Ala	GCC Ala	GCC Ala 345	GAG Glu	AGC Ser	GAG Glu	CGC Arg	TTC Phe 350	GTC Val	CGG Arg	1056
	30	CAG Gln	GGC Gly	ACC Thr 355	GGC Gly	AAC Asn	GAC Asp	GAG Glu	GCC Ala 360	GGC Gly	GCG Ala	GCC Ala	AAC Asn	GCC Ala 365	GAC Asp	GTG Val	GTG Val	1104
	35	AGC Ser	CTG Leu 370	ACC Thr	TGC Cys	CCG Pro	GTC Val	GCC Ala 375	GCC Ala	GGT Gly	GAA Glu	TGC Cys	GCG Ala 380	GGC Gly	CCG Pro	GCG Ala	GAC Asp	1152
Juzzo	40	AGC Ser 385	GGC Gly	GAC Asp	GCC Ala	CTG Leu	CTG Leu 390	GAG Glu	CGC Arg	AAC Asn	TAT Tyr	CCC Pro 395	ACT Thr	GGC Gly	GCG Ala	GAG Glu	TTC Phe 400	1200
		CTC Leu	GGC Gly	GAC Asp	GGC Gly	GGC Gly 405	GAC Asp	GTC Val	AGC Ser	TTC Phe	AGC Ser 410	ACC Thr	CGC Arg	GGC Gly	ACG Thr	CAG Gln 415	AAC Asn	1248
	45	TGG Trp	ACG Thr	GTG Val	GAG Glu 420	CGG Arg	CTG Leu	CTC Leu	CAG Gln	GCG Ala 425	CAC His	CGC Arg	CAA Gln	Leu	GAG Glu 430	GIu	CGC Arg	1296
	50	GGC Gly	TAT Tyr	GTG Val 435	TTC Phe	GTC Val	GGC Gly	TAC Tyr	CAC His 440	GGC Gly	ACC Thr	TTC Phe	CTC Leu	GAA Glu 445	GCG Ala	GCG Ala	CAA Gln	1344
	55	AGC Ser	ATC Ile 450	GTC Val	TTC Phe	GGC Gly	GGG Gly	GTG Val 455	CGC Arg	GCG Ala	CGC Arg	AGC Ser	CAG Gln 460	GAC Asp	CTC Leu	GAC Asp	GCG Ala	1392
	60	ATC Ile 465	TGG Trp	CGC Arg	GGT Gly	TTC Phe	TAT Tyr 470	ATC Ile	GCC Ala	GGC Gly	GAT Asp	CCG Pro 475	GCG Ala	CTG Leu	GCC Ala	TAC Tyr	GGC Gly 480	1440
	<i></i>	TAC Tyr	GCC Ala	CAG Gln	GAC Asp	CAG Gln 485	GAA Glu	CCC Pro	GAC Asp	GCA Ala	CGC Arg 490	GGC Gly	CGG Arg	ATC Ile	CGC Arg	AAC Asn 495	GGT Gly	1488
	65	GCC Ala	CTG Leu	CTG Leu	CGG Arg 500	GTC Val	TAT Tyr	GTG Val	CCG Pro	CGC Arg 505	TCG Ser	AGC Ser	CTG Leu	CCG Pro	GGC Gly 510	TTC Phe	TAC Tyr	1536



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	CGC Arg	ACC Thr	AGC Ser 515	CTG Leu	ACC Thr	CTG Leu	GCC Ala	GCG Ala 520	CCG Pro	GAG Glu	GCG Ala	GCG Ala	GGC Gly 525	GAG Glu	GTC Val	GAA Glu	1584
5	CGG Arg	CTG Leu 530	ATC Ile	GGC Gly	CAT His	CCG Pro	CTG Leu 535	CCG Pro	CTG Leu	CGC Arg	CTG Leu	GAC Asp 540	GCC Ala	ATC Ile	ACC Thr	GGC Gly	1632
10	CCC Pro 545	GAG Glu	GAG Glu	GAA Glu	GGC Gly	GGG Gly 550	CGC Arg	CTG Leu	GAG Glu	ACC Thr	ATT Ile 555	CTC Leu	GGC Gly	TGG Trp	CCG Pro	CTG Leu 560	1680
15	GCC Ala	GAG Glu	CGC Arg	ACC Thr	GTG Val 565	GTG Val	ATT Ile	CCC Pro	TCG Ser	GCG Ala 570	ATC Ile	CCC Pro	ACC Thr	GAC Asp	CCG Pro 575	CGC Arg	1728
	AAC Asn	GTC Val	GGC Gly	GGC Gly 580	GAC Asp	CTC Leu	GAC Asp	CCG Pro	TCC Ser 585	AGC Ser	ATC Ile	CCC Pro	GAC Asp	AAG Lys 590	GAA Glu	CAG Gln	1776
20	GCG Ala	ATC Ile	AGC Ser 595	GCC Ala	CTG Leu	CCG Pro	GAC Asp	TAC Tyr 600	GCC Ala	AGC Ser	CAG Gln	CCC Pro	GGC Gly 605	AAA Lys	CCG Pro	CCG Pro	1824
25				CTG Leu													1839
30																	